

## SEQUENCE LISTING

<110> Boehringer Ingelheim International GmbH

<120> Method for identifying compounds that modulate sister chromatid separation

<130> 0652.2290001

<140> To be assigned

<141> Herewith

<150> EP 01 101 252.3

<151> 2001-01-19

<150> US 60/297,440

<151> 2001-06-13

<160> 8

<170> PatentIn Ver. 2.1

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<211> 6668

<212> DNA

<213> Homo sapiens

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Lys Ala Gln Lys His Arg Gly Ser Leu Glu Ile Ala Asp Gln Leu Gln	
1645 1650 1655	
ggg ctg agc ctt cag gag atg cct gga gat gtc ccc ctg gcc cgc atc	5163
Gly Leu Ser Leu Gln Glu Met Pro Gly Asp Val Pro Leu Ala Arg Ile	
1660 1665 1670	
cag cgc ctc ttt tcc ttc agg gct ttg gaa tct ggc cac ttc ccc cag	5211
Gln Arg Leu Phe Ser Phe Arg Ala Leu Glu Ser Gly His Phe Pro Gln	
1675 1680 1685	
cct gaa aag gag agt ttc cag gag cgc ctg gct ctg atc ccc agt ggg	5259
Pro Glu Lys Glu Ser Phe Gln Glu Arg Leu Ala Leu Ile Pro Ser Gly	
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Val Thr Val Cys Val Leu Ala Leu Ala Thr Leu Gln Pro Gly Thr Val	
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Gly Asn Thr Leu Leu Thr Arg Leu Glu Lys Asp Ser Pro Pro Val	
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agt gtg cag att ccc act ggc cag aac aag ctt cat ctg cgt tca gtc	5403
Ser Val Gln Ile Pro Thr Gly Gln Asn Lys Leu His Leu Arg Ser Val	
1740 1745 1750	
ctg aat gag ttt gat gcc atc cag aag gca cag aaa gag aac agc agc	5451
Leu Asn Glu Phe Asp Ala Ile Gln Lys Ala Gln Lys Glu Asn Ser Ser	
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Cys Thr Asp Lys Arg Glu Trp Trp Thr Gly Arg Leu Ala Leu Asp His	
1770 1775 1780 1785	
agg atg gag gtt ctc atc gct tcc cta gag aag tct gtg ctg ggc tgc	5547
Arg Met Glu Val Leu Ile Ala Ser Leu Glu Lys Ser Val Leu Gly Cys	
1790 1795 1800	
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Trp Lys Gly Leu Leu Leu Pro Ser Ser Glu Glu Pro Gly Pro Ala Gln	

1805	1810	1815	
gag gcc tcc cgc cta cag Glu Ala Ser Arg Leu Gln 1820	gag ctg cta cag gac Glu Leu Leu Gln Asp 1825	tgt ggc tgg aaa tat Cys Gly Trp Lys Tyr 1830	5643
cct gac cgc act ctg ctg Pro Asp Arg Thr Leu Leu 1835	aaa atc atg ctc agt ggt Lys Ile Met Leu Ser Gly 1840	gcc ggt gcc ctc Ala Gly Ala Leu 1845	5691
acc cct cag gac att cag Thr Pro Gln Asp Ile Gln 1850	gcc ctg gcc tac ggg ctg Ala Leu Ala Tyr Gly Leu 1855	tgc cca acc cag Cys Pro Thr Gln 1860	5739
cca gag cga gcc cag gag Pro Glu Arg Ala Gln Glu 1870	ctc ctg aat gag gca gta Leu Leu Asn Glu Ala Val 1875	gga cgt cta cag Gly Arg Leu Gln 1880	5787
ggc ctg aca gta cca agc Gly Leu Thr Val Pro Ser 1885	aat agc cac ctt gtc ttg Asn Ser His Leu Val Leu 1890	gtc cta gac aag Val Leu Asp Lys 1895	5835
gac ttg cag aag ctg ccg Asp Leu Gln Lys Leu Pro 1900	tgg gaa agc atg ccc agc Trp Glu Ser Met Pro Ser 1905	ctc caa gca ctg Leu Gln Ala Leu 1910	5883
cct gtc acc cgg ctg ccc Pro Val Thr Arg Leu Pro 1915	tcc ttc cgc ttc cta ctc Ser Phe Arg Phe Leu Leu 1920	agc tac tcc atc Ser Tyr Ser Ile 1925	5931
atc aaa gag tat ggg gcc Ile Lys Glu Tyr Gly Ala 1930	tcg cca gtg ctg agt caa Ser Pro Val Leu Ser Gln 1935	ggg gtg gat cca Gly Val Asp Pro 1940	5979
cga agt acc ttc tat gtc Arg Ser Thr Phe Tyr Val 1950	ctg aac cct cac aat aac Leu Asn Pro His Asn Asn 1955	ctg tca agc aca Leu Ser Ser Thr 1960	6027
gag gag caa ttt cga gcc Glu Glu Gln Phe Arg Ala 1965	aat ttc agc agt gaa gct Asn Phe Ser Ser Glu Ala 1970	ggc tgg aga gga Gly Trp Arg Gly 1975	6075
gtg gtt ggg gag gtg cca Val Val Gly Glu Val Pro 1980	aga cct gaa cag gtg cag Arg Pro Glu Gln Val Gln 1985	gaa gcc ctg aca Glu Ala Leu Thr 1990	6123
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ctt gat ggg cag gct gtc Leu Asp Gly Gln Ala Val 2010	ctg cgg ctg agc tgt cgg Leu Arg Leu Ser Cys Arg 2015	gca gtg gcc ctg Ala Val Ala Leu 2020	6219
ctg ttt ggc tgt agc agt Leu Phe Gly Cys Ser Ser 2030	gcg gcc ctg gct gtg cat Ala Ala Leu Ala Val His 2035	gga aac ctg gag Gly Asn Leu Glu 2040	6267
ggg gct ggc atc gtg ctc Gly Ala Gly Ile Val Leu 2045	aag tac atc atg gct ggt Lys Tyr Ile Met Ala Gly 2050	tgc ccc ttg ttt Cys Pro Leu Phe 2055	6315

ctg ggt aat ctc tgg gat gtg act gac cgc gac att gac cgc tac acg 6363  
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Pro Pro Ala Gly Phe Pro Ser Ser Arg Ser Asp Ala Glu Arg Arg Gln  
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Ala Cys Asp Ala Ile Leu Arg Ala Cys Asn Gln Gln Leu Thr Ala Lys  
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Leu Ala Cys Pro Arg His Leu Gly Ser Leu Leu Glu Leu Ala Glu Leu  
 65 70 75 80

Ala Cys Asp Gly Tyr Leu Val Ser Thr Pro Gln Arg Pro Pro Leu Tyr  
 85 90 95

Leu Glu Arg Ile Leu Phe Val Leu Leu Arg Asn Ala Ala Ala Gln Gly  
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Ser Pro Glu Val Thr Leu Arg Leu Ala Gln Pro Leu His Ala Cys Leu  
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Val Gln Cys Ser Arg Glu Ala Ala Pro Gln Asp Tyr Glu Ala Val Ala  
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Arg Gly Ser Phe Ser Leu Leu Trp Lys Gly Ala Glu Ala Leu Leu Glu  
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Arg Arg Ala Ala Phe Ala Ala Arg Leu Lys Ala Leu Ser Phe Leu Val  
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Pro	Thr	Ala	Cys	Arg	Ala	Val	Ala	Ala	His	Gln	Leu	Phe	Asp	Ala	Ser			
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Glu	His	Cys	Arg	Arg	Phe	Cys	Trp	Ser	Arg	His	His	Asp	Lys	Ala	Ile			
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Ser	Ala	Val	Glu	Lys	Ala	His	Ser	Tyr	Leu	Arg	Asn	Thr	Asn	Leu	Ala			
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Pro	Ser	Leu	Gln	Leu	Cys	Gln	Leu	Gly	Val	Lys	Leu	Leu	Gln	Val	Gly			
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Glu	Glu	Gly	Pro	Gln	Ala	Val	Ala	Lys	Leu	Leu	Ile	Lys	Ala	Ser	Ala			
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Val	Leu	Ser	Lys	Ser	Met	Glu	Ala	Pro	Ser	Pro	Pro	Leu	Arg	Ala	Leu			
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Tyr	Glu	Ser	Cys	Gln	Phe	Phe	Leu	Ser	Gly	Leu	Glu	Arg	Gly	Thr	Lys			
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Ile	Val	Asp	Leu	Ala	Asp	Leu	Thr	Gln	Leu	Val	Asp	Ser	Cys	Lys	Ser			
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Thr	Val	Val	Trp	Met	Leu	Glu	Ala	Leu	Glu	Gly	Leu	Ser	Gly	Gln	Glu			
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 Leu Trp Leu Ala Ala Leu Gln Pro Cys Ser Pro Glu His Met Ala Glu  
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 Pro Val Thr Phe Trp Val Arg Val Lys Met Asp Ala Ala Arg Ala Gly  
 545 550 555 560  
 Asp Lys Glu Leu Gln Leu Lys Thr Leu Arg Asp Ser Leu Ser Gly Trp  
 565 570 575  
 Asp Pro Glu Thr Leu Ala Leu Leu Leu Arg Glu Glu Leu Gln Ala Tyr  
 580 585 590  
 Lys Ala Val Arg Ala Asp Thr Gly Gln Glu Arg Phe Asn Ile Ile Cys  
 595 600 605  
 Asp Leu Leu Glu Leu Ser Pro Glu Glu Thr Pro Ala Gly Ala Trp Ala  
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 625 630 635 640  
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 645 650 655  
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 675 680 685  
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 690 695 700  
 Ala Pro Gly Asn Leu Glu Glu Phe Glu Val Asn Asp Leu Asn Tyr Glu  
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 Asp Lys Leu Gln Glu Asp Arg Phe Leu Tyr Ser Asn Ile Ala Phe Asn  
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 Cys His Ile Thr Gln Leu Leu Leu Thr Leu Gly Cys Pro Ser Tyr Ala  
 820 825 830  
 Gln Leu His Leu Glu Glu Ala Ala Ser Ser Leu Lys His Leu Asp Gln

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Thr	Gln	Lys	Ala	Ala	Val	Glu	Thr	Ser	Phe	Leu	Asp	Tyr	Gly	Glu	Asn
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Pro	Asn	Phe	Leu	Ser	His	Ser	Pro	Thr	Cys	Asp	Cys	Ser	Leu	Cys	Ala
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 Pro Glu Glu Glu Leu Thr Asp Asn Trp Arg Lys Met Ser Phe Glu Ile  
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Ile Met Leu Ser Gly Ala Gly Ala Leu Thr Pro Gln Asp Ile Gln Ala  
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 Ser His Leu Val Leu Val Leu Asp Lys Asp Leu Gln Lys Leu Pro Trp  
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 Glu Ser Met Pro Ser Leu Gln Ala Leu Pro Val Thr Arg Leu Pro Ser  
 1905 1910 1915 1920  
 Phe Arg Phe Leu Leu Ser Tyr Ser Ile Ile Lys Glu Tyr Gly Ala Ser  
 1925 1930 1935  
 Pro Val Leu Ser Gln Gly Val Asp Pro Arg Ser Thr Phe Tyr Val Leu  
 1940 1945 1950  
 Asn Pro His Asn Asn Leu Ser Ser Thr Glu Glu Gln Phe Arg Ala Asn  
 1955 1960 1965  
 Phe Ser Ser Glu Ala Gly Trp Arg Gly Val Val Gly Glu Val Pro Arg  
 1970 1975 1980  
 Pro Glu Gln Val Gln Glu Ala Leu Thr Lys His Asp Leu Tyr Ile Tyr  
 1985 1990 1995 2000  
 Ala Gly His Gly Ala Gly Ala Arg Phe Leu Asp Gly Gln Ala Val Leu  
 2005 2010 2015  
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Tyr Gly Leu Pro Val Ser Leu Arg  
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